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RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/389,000

BATCH (1643)  
(3-10)

DATE: 03/22/2000  
TIME: 11:54:22

Input Set: I389000.RAW

This Raw Listing contains the General Information  
Section and up to first 5 pages.

ENTERED

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1 <110> APPLICANT: Afar, Daniel E.
2   Hubert, Rene S.
3   Raitano, Arthur B.
4 <120> TITLE OF INVENTION: PHELI: A TESTIS-SPECIFIC PROTEIN EXPRESSED IN CANCER
5 <130> FILE REFERENCE: 1703-018.US1
6 <140> CURRENT APPLICATION NUMBER: US/09/389,000
7 <141> CURRENT FILING DATE: 1999-08-31
8 <150> EARLIER APPLICATION NUMBER: 60/098,610
9 <151> EARLIER FILING DATE: 1998-08-31
10 <150> EARLIER APPLICATION NUMBER: 60/106,524
11 <151> EARLIER FILING DATE: 1998-10-03
12 <160> NUMBER OF SEQ ID NOS: 12
13 <170> SOFTWARE: PatentIn Ver. 2.1
14 <210> SEQ ID NO 1
15 <211> LENGTH: 2128
16 <212> TYPE: DNA
17 <213> ORGANISM: Homo sapiens
18 <400> SEQUENCE: 1
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20   tgtcttgtgc tctaaggtgc tgaggggaaa gacgcgggag gtctctggcc tgacactatg 120
21   aaggaagaga gaaactacaa cttcgacggt gtgagcacca accgcctgaa acagcagttg 180
22   ctggaagaag tccgcaagaa gtagtgaatg gaaaacccgt tatgagacac aacttgaatt 240
23   aaatgatgaa ctagaaaagc aaattgttta tctcaaggag aaagtggaaa aaatccatgg 300
24   aaactcttca gatagactat cttctattcg tgtctatgaa cgaatgccag tggaatcctt 360
25   aaacacatta cttaaacagc tagaagaaga aaagaagact cttgaaagtc aagtgaataa 420
26   ctatgcactt aaactggaac aagaatcaaa ggcttaccag aagatcaaca atgaacgccg 480
27   tacataccta gctgaaatgt ctcaggggtc tgggtttacat caagtttcta aaaggcaaca 540
28   ggtggatcaa ctgcctagga tgcaagagaa tctagtgaaa acgcaaaaat agacatctta 600
29   ttagttggag atgtcactgt gggctacctg gctgatactg tacagaaact atttgcaaac 660
30   atagcagaag tcaccatcac catcagtgac acgaaggagg cagcagcgct tttggatgat 720
31   tgcataattca acatggttct cttgaagggt ccttcttcac taagtgccga ggagctggaa 780
32   gccatcaagt taattagatt tggcaaaaag aaaaatacac attcactgtt tgtttttata 840
33   atccctgaaa attttaaagg ttgtatttca gggcatggaa tggatattgc ttttaactgaa 900
34   ccactgacaa tggaaaaaat gagtaatgtg gtaaaatact ggacaacatg tccctcaaac 960
35   actgttaaga ctgaaaacgc aactgggcct gaagaacttg gattgcccct gcagaggtcc 1020
36   tacagcgaac acctgggata ttttcctact gatctatttg cctgctctga atctttaagg 1080
37   aatggcaatg ggcttgaatt aaatgcttcg ttgtcagagt tcgagaaaaa caaaaagatc 1140
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39   cagctgcgta ctctcttgcc gtatgtaaaa gggagaaaaga atgatgcggc ttcagttctt 1260
40   gaggcaacag ttgattatgt gaaatatatc cgggagaaaa tctctccagc cgttatggcc 1320
41   cagattacag aagcacttca gagcaacatg aggttttgta agaaacaaca aacacccatt 1380
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43   tcccctgaga gagggctcca attcctgact aatacgtgct ggaatgggtg ctccactcct 1500
44   gatgcagaga gtccttggga tgaagctgtg agagttccat caagctccgc ctgagagaat 1560
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45      gctattggtg atccatataa aactcacatt tccagtgcag cgctgtctct gaattccttg 1620
46      catactgtca gatattattc taaagtcacc ccttcctacg atgcaactgc tgtaacaaat 1680
47      cagaacattt caattcattt accttcagcc atgcccccg tctcaagctt ctccctcggc 1740
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49      gggcgtatta aaagcacatg tttgaaattc acactctcaa ccacctactg ggcgcagttt 1860
50      gacaatctag gaaaagtgga acaaaagaatg attttgaaag ctccacccaa agacctata 1920
51      tcaaaagagt tggcatggtt tggcttctga taaatgcact caaagcttct gcagatagaa 1980
52      agaccagcag cgaaaaagct ggccacacac tgtcactcat cttcatacac acttggatcc 2040
53      ccgccagcca gagagctaca agaacaaatg gcctcagtga cctacactct ctttttctcaa 2100
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55 &lt;210&gt; SEQ ID NO 2

56 &lt;211&gt; LENGTH: 405

57 &lt;212&gt; TYPE: PRT

58 &lt;213&gt; ORGANISM: Homo sapiens

59 &lt;400&gt; SEQUENCE: 2

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60      Met Val Leu Leu Lys Val Pro Ser Ser Leu Ser Ala Glu Glu Leu Glu
61      1          5          10          15
62      Ala Ile Lys Leu Ile Arg Phe Gly Lys Lys Lys Asn Thr His Ser Leu
63      20          25          30
64      Phe Val Phe Ile Ile Pro Glu Asn Phe Lys Gly Cys Ile Ser Gly His
65      35          40          45
66      Gly Met Asp Ile Ala Leu Thr Glu Pro Leu Thr Met Glu Lys Met Ser
67      50          55          60
68      Asn Val Val Lys Tyr Trp Thr Thr Cys Pro Ser Asn Thr Val Lys Thr
69      65          70          75          80
70      Glu Asn Ala Thr Gly Pro Glu Glu Leu Gly Leu Pro Leu Gln Arg Ser
71      85          90          95
72      Tyr Ser Glu His Leu Gly Tyr Phe Pro Thr Asp Leu Phe Ala Cys Ser
73      100          105          110
74      Glu Ser Leu Arg Asn Gly Asn Gly Leu Glu Leu Asn Ala Ser Leu Ser
75      115          120          125
76      Glu Phe Glu Lys Asn Lys Lys Ile Ser Leu Leu His Ser Ser Lys Glu
77      130          135          140
78      Lys Leu Arg Arg Glu Arg Ile Lys Tyr Cys Cys Glu Gln Leu Arg Thr
79      145          150          155          160
80      Leu Leu Pro Tyr Val Lys Gly Arg Lys Asn Asp Ala Ala Ser Val Leu
81      165          170          175
82      Glu Ala Thr Val Asp Tyr Val Lys Tyr Ile Arg Glu Lys Ile Ser Pro
83      180          185          190
84      Ala Val Met Ala Gln Ile Thr Glu Ala Leu Gln Ser Asn Met Arg Phe
85      195          200          205
86      Cys Lys Lys Gln Gln Thr Pro Ile Glu Leu Ser Leu Pro Gly Thr Val
87      210          215          220
88      Met Ala Gln Arg Glu Asn Ser Val Met Ser Thr Tyr Ser Pro Glu Arg
89      225          230          235          240
90      Gly Leu Gln Phe Leu Thr Asn Thr Cys Trp Asn Gly Cys Ser Thr Pro
91      245          250          255
92      Asp Ala Glu Ser Ser Leu Asp Glu Ala Val Arg Val Pro Ser Ser Ser
93      260          265          270
94      Ala Ser Glu Asn Ala Ile Gly Asp Pro Tyr Lys Thr His Ile Ser Ser

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95          275          280          285
96      Ala Ala Leu Ser Leu Asn Ser Leu His Thr Val Arg Tyr Tyr Ser Lys
97          290          295          300
98      Val Thr Pro Ser Tyr Asp Ala Thr Ala Val Thr Asn Gln Asn Ile Ser
99      305          310          315          320
100     Ile His Leu Pro Ser Ala Met Pro Pro Val Ser Ser Phe Ser Leu Gly
101          325          330          335
102     Thr Ala Leu Leu Gly Trp Ala Arg Arg Ala Leu His Ile Pro Thr Val
103          340          345          350
104     Cys Asn Ser Phe Gly Arg Ile Lys Ser Thr Cys Leu Lys Phe Thr Leu
105          355          360          365
106     Ser Thr Thr Tyr Trp Ala Gln Phe Asp Asn Leu Gly Lys Val Glu Gln
107          370          375          380
108     Arg Met Ile Leu Lys Ala Pro Pro Lys Asp Leu Ile Ser Lys Glu Leu
109     385          390          395          400
110     Ala Trp Phe Gly Phe
111          405

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112 <210> SEQ ID NO 3
113 <211> LENGTH: 50
114 <212> TYPE: PRT
115 <213> ORGANISM: Rattus norvegicus
116 <400> SEQUENCE: 3
117     His Asn Ala Leu Glu Arg Lys Arg Arg Asp His Ile Lys Asp Ser Phe
118         1          5          10          15
119     His Ser Leu Arg Asp Ser Val Pro Ser Leu Gln Gly Glu Lys Ala Ser
120         20          25          30
121     Arg Ala Gln Ile Leu Asp Lys Ala Thr Glu Tyr Ile Gln Tyr Met Arg
122         35          40          45
123     Arg Lys
124         50

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125 <210> SEQ ID NO 4
126 <211> LENGTH: 24
127 <212> TYPE: PRT
128 <213> ORGANISM: Brachydanio rerio
129 <400> SEQUENCE: 4
130     His Asn Glu Leu Glu Lys Asn Arg Arg Ala His Leu Arg Leu Cys Leu
131         1          5          10          15
132     Glu Arg Leu Lys Thr Leu Ile Pro
133         20

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134 <210> SEQ ID NO 5
135 <211> LENGTH: 14
136 <212> TYPE: DNA
137 <213> ORGANISM: Artificial Sequence
138 <220> FEATURE:
139 <223> OTHER INFORMATION: Description of Artificial Sequence: cDNA synthesis
140     primer
141 <400> SEQUENCE: 5
142     ttttgatcaa gctt
143 <210> SEQ ID NO 6
144 <211> LENGTH: 42

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146 <213> ORGANISM: Artificial Sequence  
147 <220> FEATURE:  
148 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Adaptor 1  
149 <400> SEQUENCE: 6  
150 ctaatacgac tcactatagg gctcgagcgg ccgcccgggc ag 42  
151 <210> SEQ ID NO 7  
152 <211> LENGTH: 40  
153 <212> TYPE: DNA  
154 <213> ORGANISM: Artificial Sequence  
155 <220> FEATURE:  
156 <223> OTHER INFORMATION: Description of Artificial Sequence: DNA Adaptor 2  
157 <400> SEQUENCE: 7  
158 gtaatacgac tcactatagg gcagcgtggt cgcgcccgag 40  
159 <210> SEQ ID NO 8  
160 <211> LENGTH: 22  
161 <212> TYPE: DNA  
162 <213> ORGANISM: Artificial Sequence  
163 <220> FEATURE:  
164 <223> OTHER INFORMATION: Description of Artificial Sequence: PCR primer 1  
165 <400> SEQUENCE: 8  
166 ctaatacgac tcactatagg gc 22  
167 <210> SEQ ID NO 9  
168 <211> LENGTH: 22  
169 <212> TYPE: DNA  
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171 <220> FEATURE:  
172 <223> OTHER INFORMATION: Description of Artificial Sequence: Nested primer  
173 (NP) 1  
174 <400> SEQUENCE: 9  
175 tcgagcggcc gcccgggcag ga 22  
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179 <213> ORGANISM: Artificial Sequence  
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184 agcgtggtcg cggccgagga 20  
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186 <211> LENGTH: 23  
187 <212> TYPE: DNA  
188 <213> ORGANISM: Artificial Sequence  
189 <220> FEATURE:  
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191 22P4G9.1  
192 <400> SEQUENCE: 11  
193 ctgcgtactc tcttgccgta tgt 23  
194 <210> SEQ ID NO 12

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196 <212> TYPE: DNA  
197 <213> ORGANISM: Artificial Sequence  
198 <220> FEATURE:  
199 <223> OTHER INFORMATION: Description of Artificial Sequence: RT-PCR primer  
200 22P4G9.2  
201 <400> SEQUENCE: 12  
202 gctcaatggg tgtttggtgt ttct

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VERIFICATION SUMMARY  
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Line ? Error/Warning

Original Text

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